



#3

## SEQUENCE LISTING

&lt;110&gt; Koide, Shohei

<120> METHOD OF IDENTIFYING POLYPEPTIDE MONOBODIES WHICH BIND  
TO TARGET PROTEINS AND USE THEREOF

&lt;130&gt; 176/60901

&lt;140&gt; 10/006,760

&lt;141&gt; 2001-11-19

&lt;150&gt; 60/249,756

&lt;151&gt; 2000-11-17

&lt;160&gt; 73

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 308

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 1

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ctgctgatca gctgggatgc tccgtcagtt accgtgcgtt attaccgtat cacgtacggt 120
gaaaccggtg gtaactcccc gggttcaggaa ttactgtac ctgggttccaa gtctactgct 180
accatcagcg gcctgaaacc gggtgtcgac tataccatca ctgtatacgc tggtactggc 240
cgtggtgaca gccacgcgag ctccaagcca atctcgatta actaccgtac ctagtaactc 300
gaggatcc                                     308
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&lt;210&gt; 2

&lt;211&gt; 96

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 2

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Met Gln Val Ser Asp Val Pro Arg Asp Leu Glu Val Val Ala Ala Thr
  1             5             10             15

Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Ala Val Thr Val Arg
      20             25             30

Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val Gln
  35             40             45
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Glu Phe Thr Val Pro Gly Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu  
 50 55 60

Lys Pro Gly Val Asp Tyr Thr Ile Thr Val Tyr Ala Val Thr Gly Arg  
 65 70 75 80

Gly Asp Ser Pro Ala Ser Ser Lys Pro Ile Ser Ile Asn Tyr Arg Thr  
 85 90 95

<210> 3

<211> 96

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: mutant tenth  
 fibronectin type 3 domain of human fibronectin

<220>

<221> UNSURE

<222> (9)

<223> X at position 9 is either Asn or Lys

<400> 3

Met Gln Val Ser Asp Val Pro Arg Xaa Leu Glu Val Val Ala Ala Thr  
 1 5 10 15

Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Ala Val Thr Val Arg  
 20 25 30

Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val Gln  
 35 40 45

Glu Phe Thr Val Pro Gly Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu  
 50 55 60

Lys Pro Gly Val Asp Tyr Thr Ile Thr Val Tyr Ala Val Thr Gly Arg  
 65 70 75 80

Gly Asp Ser Pro Ala Ser Ser Lys Pro Ile Ser Ile Asn Tyr Arg Thr  
 85 90 95

<210> 4  
<211> 618  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: B42-FNfn10  
fusion protein coding region

<220>  
<221> unsure  
<222> (112)..(113)  
<223> N at positions 112 and 113 can be A, C, T, or G

<220>  
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<222> (115)..(116)  
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<220>  
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<223> N at positions 271 and 272 can be A, C, T, or G

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<221> unsure

<222> (274)..(275)  
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 <222> (280)..(281)  
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 <223> N at positions 283 and 284 can be A, C, T, or G  
  
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 <222> (286)..(287)  
 <223> N at positions 286 and 287 can be A, C, T, or G  
  
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 <222> (114)  
 <223> K at position 114 can be G or T  
  
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 <221> unsure  
 <222> (117)  
 <223> K at position 117 can be G or T  
  
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 <222> (120)  
 <223> K at position 120 can be G or T  
  
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<220>  
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 <223> K at position 273 can be G or T

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 <223> K at position 285 can be G or T

<220>  
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 <222> (288)  
 <223> K at position 288 can be G or T

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 nnknnktatt accgtatcac gtacggtgaa accggtggta actccccggt tcaggaattc 180  
 actgtacctg gttccaagtc tactgctacc atcagcggcc tgaaaccggg tgtcgactat 240  
 accatcactg tatacgtgtt tactggcnnk nnknnknnkn nknnknnktc caagccaatc 300  
 tcgattaact accgtaccag tggtagcggg ggttccccctc caaaaaagaa gagaaaggta 360  
 gctggtatca ataaagatat cgaggagtgc aatgccatca ttgagcagtt tatcgactac 420  
 ctgcgcaccg gacaggagat gccgatggaa atggcggatc aggcgattaa cgtggtgccg 480  
 ggcatgacgc cgaaaaccat tcttcacgcc gggccgccga tccagcctga ctggctgaaa 540  
 tcgaatggtt ttcataaat tgaagcggat gttaacgata ccagcctctt gctgagtgga 600  
 gattaactcg aggcattgc 618

<210> 5  
 <211> 201  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: artificial  
 B42-FNfn10 fusion protein

<220>  
 <221> UNSURE  
 <222> (38)..(42)  
 <223> Xaa at any position can be any amino acid

<220>  
 <221> UNSURE  
 <222> (90)..(96)  
 <223> Xaa at any position can be any amino acid

<400> 5  
 Met Asp Tyr Lys Asp Asp Asp Asp Lys Gly Met Gln Val Ser Asp Val  
 1 5 10 15  
 Pro Thr Asp Leu Glu Val Val Ala Ala Thr Pro Thr Ser Leu Leu Ile  
 20 25 30  
 Ser Trp Asp Ala Pro Xaa Xaa Xaa Xaa Xaa Tyr Tyr Arg Ile Thr Tyr  
 35 40 45  
 Gly Glu Thr Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Pro Gly  
 50 55 60  
 Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr  
 65 70 75 80  
 Thr Ile Thr Val Tyr Ala Val Thr Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa  
 85 90 95  
 Ser Lys Pro Ile Ser Ile Asn Tyr Arg Thr Ser Gly Thr Gly Gly Ser  
 100 105 110  
 Pro Pro Lys Lys Lys Arg Lys Val Ala Gly Ile Asn Lys Asp Ile Glu  
 115 120 125  
 Glu Cys Asn Ala Ile Ile Glu Gln Phe Ile Asp Tyr Leu Arg Thr Gly  
 130 135 140

Gln Glu Met Pro Met Glu Met Ala Asp Gln Ala Ile Asn Val Val Pro  
145 150 155 160

Gly Met Thr Pro Lys Thr Ile Leu His Ala Gly Pro Pro Ile Gln Pro  
165 170 175

Asp Trp Leu Lys Ser Asn Gly Phe His Glu Ile Glu Ala Asp Val Asn  
180 185 190

Asp Thr Ser Leu Leu Leu Ser Gly Asp  
195 200

<210> 6

<211> 96

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FNfn10  
polypeptide monobody

<220>

<221> UNSURE

<222> (28)..(32)

<223> Xaa at any position can be any amino acid

<220>

<221> UNSURE

<222> (80)..(86)

<223> Xaa at any position can be any amino acid

<400> 6

Met Gln Val Ser Asp Val Pro Thr Asp Leu Glu Val Val Ala Ala Thr  
1 5 10 15

Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Xaa Xaa Xaa Xaa Xaa  
20 25 30

Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val Gln  
35 40 45

Glu Phe Thr Val Pro Gly Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu  
50 55 60

Lys Pro Gly Val Asp Tyr Thr Ile Thr Val Tyr Ala Val Thr Gly Xaa  
65 70 75 80





<222> (427)..(428)

<223> N at positions 427 and 428 can be A, C, T, or G

<220>

<221> unsure

<222> (411)

<223> K at position 411 can be G or C

<220>

<221> unsure

<222> (414)

<223> K at position 414 can be G or C

<220>

<221> unsure

<222> (417)

<223> K at position 417 can be G or C

<220>

<221> unsure

<222> (420)

<223> K at position 420 can be G or C

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<221> unsure

<222> (423)

<223> K at position 423 can be G or C

<220>

<221> unsure

<222> (426)

<223> K at position 426 can be G or C

<220>

<221> unsure

<222> (428)

<223> K at position 428 can be G or C

<400> 7

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cctccaaaaa agaagagaaa ggtagctggt atcaataaag atatcgagga gtgcaatgcc 120  
atcattgagc agtttatcga ctacctgcgc accggacagg agatgccgat ggaaatggcg 180  
gatcaggcga ttaacgtggt gccgggcatg acgccgaaaa ccattcttca cgccggggccg 240  
ccgatccagc ctgactggct gaaatcgaat ggttttcatg aaattgaagc ggatgttaac 300  
gataccagcc tcttgctgag tggagatgcc tccaagcttg gtaccgagct cggatctatg 360  
caggtttctg atgttccgac cgacctggaa gttgttgctg cgaccccgnn snnnsnnsns 420  
nnsnnsnnsa ctagcctgct gatcagctgg gatgctcctg cagttaccgt gcgttattac 480  
cgtatcacgt acggtgaaac cggtggtaac tccccggttc aggaattcac tgtacctggt 540

tccaagtcta ctgctaccat cagcggcctg aaaccgggtg tcgactatac catcactgta 600  
 tacgctgtta ctggccgtgg tgacagccca gcgagctcca agccaatctc gattaactac 660  
 cgtacctagt aactcgaggc atgc 684

<210> 8

<211> 222

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: B42-FNfn10  
 fusion protein

<220>

<221> UNSURE

<222> (137)..(143)

<223> Xaa at any position can be any amino acid

<400> 8

Met Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu Asp Ser Thr Gln  
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Ala Met Gly Ala Pro Pro Lys Lys Lys Arg Lys Val Ala Gly Ile Asn  
 20 25 30

Lys Asp Ile Glu Glu Cys Asn Ala Ile Ile Glu Gln Phe Ile Asp Tyr  
 35 40 45

Leu Arg Thr Gly Gln Glu Met Pro Met Glu Met Ala Asp Gln Ala Ile  
 50 55 60

Asn Val Val Pro Gly Met Thr Pro Lys Thr Ile Leu His Ala Gly Pro  
 65 70 75 80

Pro Ile Gln Pro Asp Trp Leu Lys Ser Asn Gly Phe His Glu Ile Glu  
 85 90 95

Ala Asp Val Asn Asp Thr Ser Leu Leu Leu Ser Gly Asp Ala Ser Lys  
 100 105 110

Leu Gly Thr Glu Leu Gly Ser Met Gln Val Ser Asp Val Pro Thr Asp  
 115 120 125

Leu Glu Val Val Ala Ala Thr Pro Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr  
 130 135 140

Ser Leu Leu Ile Ser Trp Asp Ala Pro Ala Val Thr Val Arg Tyr Tyr

145	150	155	160
Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val Gln Glu Phe			
165	170	175	
Thr Val Pro Gly Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu Lys Pro			
180	185	190	
Gly Val Asp Tyr Thr Ile Thr Val Tyr Ala Val Thr Gly Arg Gly Asp			
195	200	205	
Ser Pro Ala Ser Ser Lys Pro Ile Ser Ile Asn Tyr Arg Thr			
210	215	220	

<210> 9  
 <211> 103  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: FNfn10  
 polypeptide monobody

<220>  
 <221> UNSURE  
 <222> (18)..(24)  
 <223> Xaa at any position can be any amino acid

<400> 9  
 Met Gln Val Ser Asp Val Pro Thr Asp Leu Glu Val Val Ala Ala Thr  
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 Pro Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Leu Leu Ile Ser Trp Asp  
 20 25 30  
 Ala Pro Ala Val Thr Val Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr  
 35 40 45  
 Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Pro Gly Ser Lys Ser  
 50 55 60  
 Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr Ile Thr  
 65 70 75 80  
 Val Tyr Ala Val Thr Gly Arg Gly Asp Ser Pro Ala Ser Ser Lys Pro  
 85 90 95

Ile Ser Ile Asn Tyr Arg Thr  
100

<210> 10  
<211> 704  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: B42-FNfn10  
fusion protein coding region

<220>  
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<222> (439)..(440)  
<223> N at positions 439 and 440 can be A, C, T, or G

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<222> (442)..(443)  
<223> N at positions 442 and 443 can be A, C, T, or G

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<222> (445)..(446)  
<223> N at positions 445 and 446 can be A, C, T, or G

<220>  
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<222> (448)..(449)  
<223> N at positions 448 and 449 can be A, C, T, or G

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<223> N at positions 451 and 452 can be A, C, T, or G

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<223> N at positions 595 and 596 can be A, C, T, or G

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<223> N at positions 598 and 599 can be A, C, T, or G

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 <223> N at positions 601 and 602 can be A, C, T, or G

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 <223> N at positions 604 and 605 can be A, C, T, or G

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<220>  
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 <222> (613)..(614)  
 <223> N at positions 613 and 614 can be A, C, T, or G

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 <222> (444)  
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<221> unsure

<222> (597)

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<220>

<221> unsure

<222> (600)

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<221> unsure

<222> (603)

<223> K at position 603 can be G or T

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<221> unsure

<222> (606)

<223> K at position 606 can be G or T

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<221> unsure

<222> (609)

<223> K at position 609 can be G or T

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<221> unsure

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<223> K at position 612 can be G or T

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<221> unsure

<222> (615)

<223> K at position 615 can be G or T

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atcattgagc agtttatcga ctacctgcgc accggacagg agatgccgat ggaaatggcg 180  
gatcaggcga ttaacgtggt gccgggcatg acgccgaaaa ccattcttca cgccgggccg 240  
ccgatccagc ctgactggct gaaatcgaat ggttttcatg aaattgaagc ggatgttaac 300  
gataccagcc tcttgctgag tggagatgcc tccaagcttg gtaccgagct cggatctatg 360  
caggtttctg atgttccgac cgacctgga gttgttgctg cgaccccgac tagcctgctg 420  
atcagctggg atgctcctnn knnknknkn nnktattacc gtatcacgta cggtgaaacc 480  
ggtggtaact ccccggttca ggaattcact gtacctggtt ccaagtctac tgctaccatc 540  
agcggcctga aaccgggtgt cgactatacc atcactgtat acgctgttac tggcnnknkn 600

nnknnknnkn nknnktccaa gccaatctcg attaactacc gtacctagta actcgaggca 660  
 tgcactaga gggccgcac atgtaattag ttatgtcacg ctta 704

<210> 11  
 <211> 215  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: B42-FNfn10  
 fusion protein

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 <222> (147)..(151)  
 <223> Xaa at positions 147, 148, 149, 150, and 151 can  
 be any amino acid

<220>  
 <221> UNSURE  
 <222> (199)..(205)  
 <223> Xaa at positions 199, 200, 201, 202, 203, 204, and  
 205 can be any amino acid

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 1 5 10 15  
 Ala Met Gly Ala Pro Pro Lys Lys Lys Arg Lys Val Ala Gly Ile Asn  
 20 25 30  
 Lys Asp Ile Glu Glu Cys Asn Ala Ile Ile Glu Gln Phe Ile Asp Tyr  
 35 40 45  
 Leu Arg Thr Gly Gln Glu Met Pro Met Glu Met Ala Asp Gln Ala Ile  
 50 55 60  
 Asn Val Val Pro Gly Met Thr Pro Lys Thr Ile Leu His Ala Gly Pro  
 65 70 75 80  
 Pro Ile Gln Pro Asp Trp Leu Lys Ser Asn Gly Phe His Glu Ile Glu  
 85 90 95  
 Ala Asp Val Asn Asp Thr Ser Leu Leu Leu Ser Gly Asp Ala Ser Lys  
 100 105 110  
 Leu Gly Thr Glu Leu Gly Ser Met Gln Val Ser Asp Val Pro Thr Asp

115	120	125
Leu Glu Val Val Ala Ala Thr Pro Thr Ser Leu Leu Ile Ser Trp Asp		
130	135	140
Ala Pro Xaa Xaa Xaa Xaa Xaa Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr		
145	150	155 160
Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Pro Gly Ser Lys Ser		
165	170	175
Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr Ile Thr		
180	185	190
Val Tyr Ala Val Thr Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ser Lys Pro		
195	200	205
Ile Ser Ile Asn Tyr Arg Thr		
210	215	

<210> 12

<211> 96

<212> PRT

<213> Artificial Sequence . . . .

<220>

<223> Description of Artificial Sequence: FNfn10  
polypeptide monobody

<220>

<221> UNSURE

<222> (28)..(32)

<223> Xaa at positions 28, 29, 30, 31, and 32 can be any  
amino acid

<220>

<221> UNSURE

<222> (80)..(85)

<223> Xaa at positions 80, 81, 82, 83, 84, and 85 can be  
any amino acid

<400> 12

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1 5 10 15

Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Xaa Xaa Xaa Xaa Xaa
20 25 30



Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val Gln  
35 40 45

Glu Phe Thr Val Pro Gly Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu  
50 55 60

Lys Pro Gly Val Asp Tyr Thr Ile Thr Val Tyr Ala Val Thr Gly Xaa  
65 70 75 80

Xaa Xaa Xaa Xaa Xaa Xaa Ser Lys Pro Ile Ser Ile Asn Tyr Arg Thr  
85 90 95

<210> 13

<211> 687

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: B42-FNfn10  
fusion protein coding region

<220>

<221> unsure

<222> (595)..(596)

<223> N at positions 595 and 596 can be A, C, T, or G

<220>

<221> unsure

<222> (598)..(599)

<223> N at positions 598 and 599 can be A, C, T, or G

<220>

<221> unsure

<222> (601)..(602)

<223> N at positions 601 and 602 can be A, C, T, or G

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<221> unsure

<222> (604)..(605)

<223> N at positions 604 and 605 can be A, C, T, or G

<220>

<221> unsure

<222> (607)..(608)  
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 <222> (610)..(611)  
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 <222> (619)..(620)  
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 <222> (622)..(623)  
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 <222> (625)..(626)  
 <223> N at positions 625 and 626 can be A, C, T, or G  
  
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 <223> N at positions 628 and 629 can be A, C, T, or G  
  
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 <222> (631)..(632)  
 <223> N at positions 631 and 632 can be A, C, T, or G  
  
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 <222> (634)..(635)  
 <223> N at positions can be 634 and 635 can be A, C, T,  
 or G

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<223> N at positions 640 and 641 can be A, C, T, or G

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<222> (609)  
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<222> (615)  
<223> K at position 615 can be G or T

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<222> (618)  
<223> K at position 618 can be G or T

<220>  
<221> unsure  
<222> (621)  
<223> K at position 621 can be G or T

<220>  
<221> unsure  
<222> (624)  
<223> K at position 624 can be G or T

<220>  
<221> unsure  
<222> (627)  
<223> K at position 627 can be G or T

<220>  
<221> unsure  
<222> (630)  
<223> K at position 630 can be G or T

<220>  
<221> unsure  
<222> (633)  
<223> K at position 633 can be G or T

<220>  
<221> unsure  
<222> (636)  
<223> K at position 636 can be G or T

<220>  
<221> unsure  
<222> (639)  
<223> K at position 639 can be G or T

<220>  
<221> unsure  
<222> (642)  
<223> K at position 642 can be G or T

<400> 13  
atgggtaagc ctatccctaa ccctctcctc ggtctcgatt ctacacaagc tatgggtgct 60  
cctccaaaâââ agaagagaaa ggtagctggt atcaataaag atatcgagga gtgcaatgcc 120  
atcattgagc agtttatcga ctacctgcgc accggacagg agatgccgat ggaaatggcg 180  
gatcaggcga ttaacgtggt gccgggcatg acgccgaaaa ccattcttca cgccgggccg 240

```

ccgatccagc ctgactggct gaaatcgaat ggttttcatg aaattgaagc ggatgttaac 300
gataaccagcc tcttgctgag tggagatgcc tccaagcttg gtaccgagct cggatctatg 360
cgtgtttctg atgttccgcg tgacctggaa gttgttgctg cgaccccgac tagcctgctg 420
atcagctggg atgctcctgc agttaccgtg cgttattacc gtatcacgta cggtgaaacc 480
ggtggtaact ccccggttca ggaattcact gtacctgggtt ccaagtctac tgctaccatc 540
agcggcctga aaccgggtgt cgactatacc atcactgtat acgctgttac tggcnnknnk 600
nnknnknnkn nknnknnknn knnknnknnk nnknnknnkn nkaagccaat ctcgattaac 660
taccgtacct agtaactcga ggcatagc                                     687

```

<210> 14

<211> 223

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: B42-FNfn10  
fusion protein

<220>

<221> UNSURE

<222> (199)..(214)

<223> Xaa at positions 199, 200, 201, 202, 203, 204,  
205, 206, 207, 208, 209, 210, 211, 212, 213, and  
214 can be any amino acid

<400> 14

```

Met Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu Asp Ser Thr Gln
  1              5              10              15

```

```

Ala Met Gly Ala Pro Pro Lys Lys Lys Arg Lys Val Ala Gly Ile Asn
      20              25              30

```

```

Lys Asp Ile Glu Glu Cys Asn Ala Ile Ile Glu Gln Phe Ile Asp Tyr
    35              40              45

```

```

Leu Arg Thr Gly Gln Glu Met Pro Met Glu Met Ala Asp Gln Ala Ile
    50              55              60

```

```

Asn Val Val Pro Gly Met Thr Pro Lys Thr Ile Leu His Ala Gly Pro
    65              70              75              80

```

```

Pro Ile Gln Pro Asp Trp Leu Lys Ser Asn Gly Phe His Glu Ile Glu
      85              90              95

```

```

Ala Asp Val Asn Asp Thr Ser Leu Leu Leu Ser Gly Asp Ala Ser Lys
    100              105              110

```

Leu Gly Thr Glu Leu Gly Ser Met Arg Val Ser Asp Val Pro Arg Asp  
115 120 125

Leu Glu Val Val Ala Ala Thr Pro Thr Ser Leu Leu Ile Ser Trp Asp  
130 135 140

Ala Pro Ala Val Thr Val Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr  
145 150 155 160

Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Pro Gly Ser Lys Ser  
165 170 175

Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr Ile Thr  
180 185 190

Val Tyr Ala Val Thr Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa  
195 200 205

Xaa Xaa Xaa Xaa Xaa Xaa Lys Pro Ile Ser Ile Asn Tyr Arg Thr  
210 215 220

<210> 15

<211> 104

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FNfn10  
polypeptide monobody

<220>

<221> UNSURE

<222> (80)..(95)

<223> Xaa at positions 80, 81, 82, 83, 84, 85, 86, 87,  
88, 89, 90, 91, 92, 93, 94, and 95 can be any  
amino acid

<400> 15

Met Arg Val Ser Asp Val Pro Arg Asp Leu Glu Val Val Ala Ala Thr  
1 5 10 15

Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Ala Val Thr Val Arg  
20 25 30

Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn Ser Pro Val Gln  
35 40 45

Glu Phe Thr Val Pro Gly Ser Lys Ser Thr Ala Thr Ile Ser Gly Leu  
 50 55 60

Lys Pro Gly Val Asp Tyr Thr Ile Thr Val Tyr Ala Val Thr Gly Xaa  
 65 70 75 80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Lys  
 85 90 95

Pro Ile Ser Ile Asn Tyr Arg Thr  
 100

<210> 16

<211> 663

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: B42-FNfn10  
 fusion protein coding region

<400> 16

atgggtaagc ctatccctaa cccctctcctc ggtctcgatt ctacacaagc tatgggtgct 60  
 cctccaaaaa agaagagaaa ggtagctggt atcaataaag atatcgagga gtgcaatgcc 120  
 atcattgagc agttttatcga ctacctgcgc accggacagg agatgccgat ggaaatggcg 180  
 gatcaggcga ttaacgtggt gccgggcatg acgccgaaaa ccattcttca cgccggggccg 240  
 ccgatccagc ctgactgggt gaaatcgaat ggttttcatg aaattgaagc ggatgttaac 300  
 gataccagcc tcttgctgag tggagatgcc tccaagcttg gtaccgagct cggatctatg 360  
 caggtttctg atgttccgac cgacctggaa gttgttgctg cgaccccgac tagcctgctg 420  
 atcagctggg atgctcctgc agttaccgtg cgttattacc gtatcacgta cggtgaaacc 480  
 ggtggtaact ccccggttca ggaattcact gtacctggtt ccaagtctac tgctaccatc 540  
 agcggcctga aaccgggtgt cgactatacc atcactgtat acgctgttac tggccgtggt 600  
 gacagcccag cgagctccaa gccaatctcg attaaactacc gtacctagta actcgaggca 660  
 tgc 663

<210> 17

<211> 215

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: B42-FNfn10  
 fusion protein

<400> 17

Met Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu Asp Ser Thr Gln

1	5	10	15
Ala Met Gly Ala Pro Pro Lys Lys Lys Arg Lys Val Ala Gly Ile Asn	20	25	30
Lys Asp Ile Glu Glu Cys Asn Ala Ile Ile Glu Gln Phe Ile Asp Tyr	35	40	45
Leu Arg Thr Gly Gln Glu Met Pro Met Glu Met Ala Asp Gln Ala Ile	50	55	60
Asn Val Val Pro Gly Met Thr Pro Lys Thr Ile Leu His Ala Gly Pro	65	70	75
Pro Ile Gln Pro Asp Trp Leu Lys Ser Asn Gly Phe His Glu Ile Glu	85	90	95
Ala Asp Val Asn Asp Thr Ser Leu Leu Leu Ser Gly Asp Ala Ser Lys	100	105	110
Leu Gly Thr Glu Leu Gly Ser Met Gln Val Ser Asp Val Pro Thr Asp	115	120	125
Leu Glu Val Val Ala Ala Thr Pro Thr Ser Leu Leu Ile Ser Trp Asp	130	135	140
Ala Pro Ala Val Thr Val Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr	145	150	155
Gly Gly Asn Ser Pro Val Gln Glu Phe Thr Val Pro Gly Ser Lys Ser	165	170	175
Thr Ala Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr Ile Thr	180	185	190
Val Tyr Ala Val Thr Gly Arg Gly Asp Ser Pro Ala Ser Ser Lys Pro	195	200	205
Ile Ser Ile Asn Tyr Arg Thr	210	215	

<210> 18

<211> 1542

<212> DNA

<213> Artificial Sequence

<220>



<223> Description of Artificial Sequence:

lexA-ER(alpha)EF fusion protein

<400> 18

```

atgaaagcgt taacggccag gcaacaagag gtgtttgata tcatccgtga tcacatcagc 60
cagacaggta tgccgccgac gcgtgccgaa atcgccgagc gtttgggggt cgttcccca 120
aacgcggctg aagaacatct gaaggcgctg gcacgcaaag gcgttattga aattgtttcc 180
ggcgcatcac gcgggattcg tctgttgacg gaagaggaag aagggttgcc gctggtaggt 240
cgtgtggctg ccggtgaacc acttctggcg caacagcata ttgaaggcca ttatcagggtc 300
gatccttcct tattcaagcc gaatgctgat ttctgctgc gcgtcagcgg gatgtcgatg 360
aaagatatcg gcattatgga tggtagcttg ctggcagtcg ataaaactca ggatgtacgt 420
aacggtcagg tcgttgtcgc acgtattgat gacgaagtta ccgttaagcg cctgaaaaaa 480
cagggaata aagtcgaact gttgccagaa aatagcgagt ttaaaccaat tgtcgtagat 540
cttcgtcagc agagcttcac cattgaaggg ctggcggttg gggttattcg caacggcgac 600
tggttggaat tcaagcttga gctcggcggc agcggtatga tcaaacgctc taagaagaac 660
agcctggcct tgtccctgac ggccgaccag atggtcagtg ccttggttga tgctgagccc 720
cccatactct attccgagta tgatcctacc agacccttca gtgaagcttc gatgatgggc 780
ttactgacca acctggcaga caggagctg gttcacatga tcaactgggc gaagagggtg 840
ccaggctttg tggatttgac cctccatgat cagggtccacc ttctagaatg tgcctggcta 900
gagatcctga tgattggtct cgtctggcgc tccatggagc acccagtga gctactgttt 960
gctcctaact tgctcttgga caggaaccag gaaaaatgtg tagagggcat ggtggagatc 1020
ttcgacatgc tgctggctac atcatctcgg ttccgcatga tgaatctgca gggagaggag 1080
tttgtgtgcc tcaaatctat tttttgctt aattctggag tgtacacatt tctgtccagc 1140
accctgaagt ctctggaaga gaaggacat atccaccgag tcctggacaa gatcacagac 1200
actttgatcc acctgatggc caaggcaggc ctgaccctgc agcagcagca ccagcggctg 1260
gcccagctcc tcctcatcct ctcccacatc aggcacatga gtaacaaagg catggagcat 1320
ctgtacagca tgaagtgcaa gaacgtggtg cccctctatg acctgctgct ggagatgctg 1380
gacgcccacc gcctacatgc gccactagc cgtggagggg catccgtgga ggagacggac 1440
caaagccact tggccactgc gggctctact tcatcgcat ccttgcaaaa gtattacatc 1500
acggggggagg cagagggttt ccctgccaca gtctgactcg ag 1542

```

<210> 19

<211> 511

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

lexA-ER(alpha)EF fusion protein

<400> 19

```

Met Lys Ala Leu Thr Ala Arg Gln Gln Glu Val Phe Asp Leu Ile Arg
  1                   5                   10                   15

```

```

Asp His Ile Ser Gln Thr Gly Met Pro Pro Thr Arg Ala Glu Ile Ala
      20                   25                   30

```

Gln	Arg	Leu	Gly	Phe	Arg	Ser	Pro	Asn	Ala	Ala	Glu	Glu	His	Leu	Lys	35	40	45	
Ala	Leu	Ala	Arg	Lys	Gly	Val	Ile	Glu	Ile	Val	Ser	Gly	Ala	Ser	Arg	50	55	60	
Gly	Ile	Arg	Leu	Leu	Gln	Glu	Glu	Glu	Glu	Gly	Leu	Pro	Leu	Val	Gly	65	70	75	80
Arg	Val	Ala	Ala	Gly	Glu	Pro	Leu	Leu	Ala	Gln	Gln	His	Ile	Glu	Gly	85	90	95	
His	Tyr	Gln	Val	Asp	Pro	Ser	Leu	Phe	Lys	Pro	Asn	Ala	Asp	Phe	Leu	100	105	110	
Leu	Arg	Val	Ser	Gly	Met	Ser	Met	Lys	Asp	Ile	Gly	Ile	Met	Asp	Gly	115	120	125	
Asp	Leu	Leu	Ala	Val	His	Lys	Thr	Gln	Asp	Val	Arg	Asn	Gly	Gln	Val	130	135	140	
Val	Val	Ala	Arg	Ile	Asp	Asp	Glu	Val	Thr	Val	Lys	Arg	Leu	Lys	Lys	145	150	155	160
Gln	Gly	Asn	Lys	Val	Glu	Leu	Leu	Pro	Glu	Asn	Ser	Glu	Phe	Lys	Pro	165	170	175	
Ile	Val	Val	Asp	Leu	Arg	Gln	Gln	Ser	Phe	Thr	Ile	Glu	Gly	Leu	Ala	180	185	190	
Val	Gly	Val	Ile	Arg	Asn	Gly	Asp	Trp	Leu	Glu	Phe	Lys	Leu	Glu	Leu	195	200	205	
Gly	Gly	Ser	Gly	Met	Ile	Lys	Arg	Ser	Lys	Lys	Asn	Ser	Leu	Ala	Leu	210	215	220	
Ser	Leu	Thr	Ala	Asp	Gln	Met	Val	Ser	Ala	Leu	Leu	Asp	Ala	Glu	Pro	225	230	235	240
Pro	Ile	Leu	Tyr	Ser	Glu	Tyr	Asp	Pro	Thr	Arg	Pro	Phe	Ser	Glu	Ala	245	250	255	
Ser	Met	Met	Gly	Leu	Leu	Thr	Asn	Leu	Ala	Asp	Arg	Glu	Leu	Val	His	260	265	270	
Met	Ile	Asn	Trp	Ala	Lys	Arg	Val	Pro	Gly	Phe	Val	Asp	Leu	Thr	Leu	275	280	285	

His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu Glu Ile Leu Met  
290 295 300

Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Val Lys Leu Leu Phe  
305 310 315 320

Ala Pro Asn Leu Leu Leu Asp Arg Asn Gln Gly Lys Cys Val Glu Gly  
325 330 335

Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser Ser Arg Phe Arg  
340 345 350

Met Met Asn Leu Gln Gly Glu Glu Phe Val Cys Leu Lys Ser Ile Ile  
355 360 365

Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser Thr Leu Lys Ser  
370 375 380

Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp Lys Ile Thr Asp  
385 390 395 400

Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr Leu Gln Gln Gln  
405 410 415

His Gln Arg Leu Ala Gln Leu Leu Leu Ile Leu Ser His Ile Arg His  
420 425 430

Met Ser Asn Lys Gly Met Glu His Leu Tyr Ser Met Lys Cys Lys Asn  
435 440 445

Val Val Pro Leu Tyr Asp Leu Leu Leu Glu Met Leu Asp Ala His Arg  
450 455 460

Leu His Ala Pro Thr Ser Arg Gly Gly Ala Ser Val Glu Glu Thr Asp  
465 470 475 480

Gln Ser His Leu Ala Thr Ala Gly Ser Thr Ser Ser His Ser Leu Gln  
485 490 495

Lys Tyr Tyr Ile Thr Gly Glu Ala Glu Gly Phe Pro Ala Thr Val  
500 505 510

<210> 20

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: protein

<220>

<221> UNSURE

<222> (2)..(3)

<223> X at any position can be any amino acid

<400> 20

Leu Xaa Xaa Leu Leu

1

5

<210> 21

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: endoplasmic  
reticulum localization signal

<400> 21

Lys Asp Glu Leu

1

<210> 22

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BC loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 22

Trp Tyr Gln Gly Arg

1

5

<210> 23

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: BC loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 23  
Pro Arg Thr Lys Gln  
1 5

<210> 24  
<211> 5  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: BC loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 24  
Val Arg Arg Pro Pro  
1 5

<210> 25  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 25  
Gly Ile Leu Glu Met Leu Gln  
1 5

<210> 26  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 26

Arg Leu Arg Ala Gln Leu Val

1

5

<210> 27

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 27

Pro Val Arg Val Leu Leu Arg

1

5

<210> 28

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 28

Arg Leu Arg Asp Leu Leu Gln

1

5

<210> 29

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 29

Gly Leu Val Ser Leu Leu Arg

1

5

<210> 30

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 30

Arg Lys Val Val Trp Thr Gly

1

5

<210> 31

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pFNB42B5F7  
library

<400> 31

Thr Ala Ala Ile Met Val Lys

1

5

<210> 32

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: consensus  
sequence

<220>

<221> UNSURE

<222> (2)..(3)

<223> X at any position can be an amino acid

<400> 32

Leu Xaa Xaa Met Leu  
1 5

<210> 33  
<211> 5  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: sequence  
within helix 12 of estrogen receptor-alpha and  
estrogen receptor-beta

<400> 33  
Leu Leu Glu Met Leu  
1 5

<210> 34  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: AB loop  
sequence for polypeptide monobody in pYT45AB7N  
library

<400> 34  
Trp Thr Trp Val Leu Arg Glu  
1 5

<210> 35  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: AB loop  
sequence for polypeptide monobody in pYT45AB7N  
library

<400> 35  
Trp Val Leu Ile Thr Arg Ser  
1 5



<210> 36  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 36  
Leu Arg Leu Met Leu Ala Gly  
1 5

<210> 37  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 37  
Ala Leu Val Glu Met Leu Arg  
1 5

<210> 38  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 38  
Arg Leu Leu Trp Asn Ser Leu  
1 5

<210> 39  
<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 39

Arg Val Leu Met Thr Leu Leu

1

5

<210> 40

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 40

Gly Leu Arg Arg Leu Leu Arg

1

5

<210> 41

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 41

Gly Leu Arg Gln Met Leu Gly

1

5

<210> 42

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 42

Arg Val Leu His Ser Leu Leu  
1 5

<210> 43

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 43

Arg Val Arg Asp Leu Leu Met  
1 5

<210> 44

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 44

Arg Val Met Asp Met Leu Leu  
1 5

<210> 45

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7

library

<400> 45

Gly Ile Ala Glu Leu Leu Arg

1

5

<210> 46

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 46

Arg Ile Leu Leu Asn Met Leu Thr

1

5

<210> 47

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 47

Gly Gly Trp Leu Trp Cys Val Thr

1

5

<210> 48

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 48

Thr Trp Val Val Arg Arg Val  
1 5

<210> 49

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 49

Thr Trp Val Arg Pro Asn Gln  
1 5

<210> 50

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 50

Arg Arg Val Pro Ile Trp Cys  
1 5

<210> 51

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 51

Arg Arg Val Tyr Asp Phe Leu  
1 5

<210> 52  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 52  
Leu Arg Gln Met Leu Ala Asp  
1 5

<210> 53  
<211> 7  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT45B3F7  
library

<400> 53  
Gly Leu Arg Met Leu Leu Arg  
1 5

<210> 54  
<211> 16  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT47F16  
library

<400> 54  
Ser Arg Arg Leu Val Glu His Leu Ala Gly Val Glu Val Gln Ala Leu  
1 5 10 15

<210> 55  
<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: FG loop  
sequence for polypeptide monobody in pYT47F16  
library

<400> 55

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<210> 64

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<400> 65  
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